SEQUENCE LISTING

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<110> Matti Sallberg
      Catharina Hultgren
<120> VACCINES CONTAINING RIBAVIRIN AND
 METHODS OF USE THEREOF
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<150> 09/705,547
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<150> 60/229,175
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Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
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Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
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Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
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Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
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Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
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Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
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Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
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Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr

185

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Asn Ser Ser Val Val Tyr Glu Ala Ala Asp Ala Ile Leu His Thr Pro
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Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ala Ser Arg Cys Trp Val
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Ala Val Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr Thr
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Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu Cys
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Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val Gly
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Gln Leu Phe Thr Phe Ser Pro Arg His His Trp Thr Thr Gln Asp Cys
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Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala Trp
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Asn Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ala Gln
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                                    330
Leu Leu Arg Ile Pro Gln Ala Ile Met Asp Met Ile Ala Gly Ala His
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Trp Gly Val Leu Ala Gly Ile Lys Tyr Phe Ser Met Val Gly Asn Trp
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Ala Lys Val Leu Val Val Leu Leu Phe Ala Gly Val Asp Ala Glu
                        375
                                            380
Thr His Val Thr Gly Gly Asn Ala Gly Arg Thr Thr Ala Gly Leu Val
                    390
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Gly Leu Leu Thr Pro Gly Ala Lys Gln Asn Ile Gln Leu Ile Asn Thr
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Asn Gly Ser Trp His Ile Asn Ser Thr Ala Leu Asn Cys Asn Glu Ser
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Leu Asn Thr Gly Trp Leu Ala Gly Leu Phe Tyr Gln His Lys Phe Asn
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Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Arg Leu Thr Asp
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Phe Ala Gln Gly Trp Gly Pro Ile Ser Tyr Ala Asn Gly Ser Gly Leu
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Asp Glu Arg Pro Tyr Cys Trp His Tyr Pro Pro Arg Pro Cys Gly Ile
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Val Pro Ala Lys Ser Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser
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Pro Val Val Val Gly Thr Thr Asp Arg Ser Gly Ala Pro Thr Tyr Ser
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Trp Gly Ala Asn Asp Thr Asp Val Phe Val Leu Asn Asn Thr Arg Pro
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Pro Leu Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Ser Thr Gly Phe
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Thr Lys Val Cys Gly Ala Pro Pro Cys Val Ile Gly Gly Val Gly Asn
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Asn Thr Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys Tyr Pro Glu Ala
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Thr Tyr Ser Arg Cys Gly Ser Gly Pro Arg Ile Thr Pro Arg Cys Met
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Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Ile Asn Tyr
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Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Leu
Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp
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Val Gly Ser Ser Ile Ala Ser Trp Ala Ile Lys Trp Glu Tyr Val Val
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Leu Leu Phe Leu Leu Ala Asp Ala Arg Val Cys Ser Cys Leu Trp
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Met Met Leu Leu Ile Ser Gln Ala Glu Ala Ala Leu Glu Asn Leu Val
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Gly Ala Val Tyr Ala Leu Tyr Gly Met Trp Pro Leu Leu Leu Leu Leu
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Leu Ala Leu Pro Gln Arg Ala Tyr Ala Leu Asp Thr Glu Val Ala Ala
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Pro Tyr Tyr Lys Arg Tyr Ile Ser Trp Cys Met Trp Trp Leu Gln Tyr
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Phe Leu Thr Arg Val Glu Ala Gln Leu His Val Trp Val Pro Pro Leu
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Asn Val Arg Gly Gly Arg Asp Ala Val Ile Leu Leu Thr Cys Val Val
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His Pro Ala Leu Val Phe Asp Ile Thr Lys Leu Leu Ala Ile Phe
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Thr Gly Thr Cys Val Tyr Asn His Leu Ala Pro Leu Arg Asp Trp Ala
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His Asn Gly Leu Arg Asp Leu Ala Val Ala Val Glu Pro Val Val Phe
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Ser Arg Met Glu Thr Lys Leu Ile Thr Trp Gly Ala Asp Thr Ala Ala
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Cys Gly Asp Ile Ile Asn Gly Leu Pro Val Ser Ala Arg Arg Gly Gln
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Glu Ile Leu Leu Gly Pro Ala Asp Gly Met Val Ser Lys Gly Trp Arg
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Leu Leu Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu
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Gly Glu Val Gln Ile Val Ser Thr Ala Thr Gln Thr Phe Leu Ala Thr
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Cys Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg
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Thr Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Thr Tyr Thr Asn Val
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    1090
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Ala Asp Val Ile Pro Val Arg Arg Gly Asp Ser Arg Gly Ser Leu 1140 1145 1150	
Leu Ser Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro 1155 1160 1165	
Leu Leu Cys Pro Thr Gly His Ala Val Gly Leu Phe Arg Ala Ala Val 1170 1175 1180	
Cys Thr Arg Gly Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Asn 1185 1190 1195 1200	
Leu Glu Thr Thr Met Arg Ser Pro Val Phe Thr Asp Asn Ser Ser Pro 1205 1210 1215	
Pro Ala Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr 1220 1225 1230	
Gly Ser Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Lys Gly 1235 1240 1245	
Tyr Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe 1250 1255 1260	
Gly Ala Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr 1265 1270 1275 1280	
Gly Val Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr 1285 1290 1295	
Gly Lys Phe Leu Ala Asp Ala Gly Cys Ser Gly Gly Ala Tyr Asp Ile 1300 1305 1310	
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Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 1330 1335 1340	
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Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val 1395 1400 1405	
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Val Ile Pro Thr Ser Gly Asp Val Val Val Ser Thr Asp Ala Leu 1425 1430 1435 1440	
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Cys Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 1460 1465 1470	
Glu Thr Thr Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg 1475 1480 1485	
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Val Arg Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln 1540 1545 1550	
Asp His Leu Gly Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile	

1555								•	
1555			560	_	_	1565			
Asp Ala His Phe 1570	Leu Ser	Gln Th	hr Lys	Gln Ser	Gly 1580		Asn	Phe	Pro
Tyr Leu Val Ala	Tyr Gln	Ala Th	hr Val	Cys Ala	Arq	Ala	Gln	Ala	Pro
1585	1590			159	_				1600
Pro Pro Ser Trp			ra Ive		_	Δra	T.e.n	Lve	
FIO FIO BEL 11p	_	INCC AL		1610	110	nr 9	LC u	1615	
Thr Leu His Gly	1605 Pro Thr	Pro Le	eu Leu	Tyr Arg	Leu	Gly	Ala		
1620			1625				1630		
Asn Glu Val Thr 1635	Leu Thr		ro Ile 640	Thr Lys	Tyr	Ile 1645		Thr	Cys
Met Ser Ala Asp	Leu Glu	Val Va	al Thr	Ser Thr	Trp	Val	Leu	Val	Glv
1650		1655			1660				1
Gly Val Leu Ala	Ala Tau		יים שניי	Cva Tax			C111	Crra	Wa I
•			ia iyi	-		1111	GIY	Cys	
1665	1670		_	167					1680
Val Ile Val Gly	Arg Ile	Val Le		Gly Lys 1690	Pro	Ala	Ile	11e 1695	
Asp Arg Glu Val	Leu Tyr	Gln Gl	lu Phe	Asp Glu	Met	Glu	Glu	Cys	Ser
1700			1705				1710		
Gln His Leu Pro 1715					Leu		Glu		Phe
	T 01			mbaa nia	0			77-	a 1
Lys Gln Lys Ala 1730	_	1735			1740)			
Val Ile Thr Pro	Ala Val	Gln Th	hr Asn	Trp Gln	Lys	Leu	Glu	Val	Phe
1745	1750			175	5				1760
Trp Ala Lys His	Met Trp	Asn Ph	he Ile	Ser Gly	Ile	Gln	Tyr	Leu	Ala
1 1	1765			1770			•	1775	
Gly Leu Ser Thr		Clv Ac			Λla	Ser	T.A11		
-		Oly As	1785		AIG	DCI	1790		niu
1780		a D-			a 1	a 1			T
Phe Thr Ala Ala 1795		18	800		_	1805			
Phe Asn Ile Leu	Glv Glv	Trp Va	al Ala	Ala Glr	Leu	בומ	Δla	Dro	Glv
1810	Cry Cry	1815	ai Aiu	mia oii	1820		nia	FIO	1
1810		1815			1820)			
	Phe Val	1815 Gly Al			1820 Gly)			
1810 Ala Ala Thr Ala 1825	Phe Val	1815 Gly Al	la Gly	Leu Ala	1820 Gly 5) Ala	Ala	Leu	Asp 1840
1810 Ala Ala Thr Ala	Phe Val 1830 Gly Lys	1815 Gly Al	la Gly	Leu Ala 183 Asp Ile	1820 Gly 5) Ala	Ala	Leu Tyr	Asp 1840 Gly
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1810 Ala Ala Thr Ala 1825 Ser Val Gly Leu Ala Gly Val Ala 1860 Val Pro Ser Thr 1875 Pro Gly Ala Leu 1890 Arg Val Gly Pro	Phe Val 1830 Gly Lys 1845 Gly Ala Clu Asp Ala Val Gly Glu 1910	1815 Gly Al Val Leu Va Leu Va Gly Va 1895 Gly Al	la Gly eu Val al Ala 1865 al Asn 880 al Val la Val	Leu Ala 183 Asp Ile 1850 Phe Lys Leu Leu Phe Ala Gln Trp	1820 Gly 5 Leu Ile Pro Ser 1900 Met	Ala Ala Met Ala 1885 Ile Asn	Ala Gly Ser 1870 Ile Leu	Leu Tyr 1855 Gly Leu Arg	Asp 1840 Gly Glu Ser Arg Ile 1920 Pro
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1810 Ala Ala Thr Ala 1825 Ser Val Gly Leu Ala Gly Val Ala 1860 Val Pro Ser Thr 1875 Pro Gly Ala Leu 1890 Arg Val Gly Pro 1905 Ala Phe Ala Ser Glu Ser Asp Ala 1940 Val Thr Gln Leu 1955 Thr Thr Pro Cys 1970	Phe Val 1830 Gly Lys 1845 Gly Ala Glu Asp Ala Val Gly Glu 1910 Arg Gly 1925 Ala Ala Leu Arg Ser Gly	1815 Gly Al Val Leu Va Leu Va 1895 Gly Al Asn Hi Arg Va Arg Le Ser Tr 1975	la Gly eu Val al Ala 1865 al Asn 880 al Val la Val is Val al Thr 1945 eu His 960 rp Leu	Leu Ala 183 Asp Ile 1850 Phe Lys Leu Leu Phe Ala Gln Trp 191 Ser Pro 1930 Ala Ile Gln Trp Arg Asp	1820 Gly 5 Leu Ile Pro Ser 1900 Met 5 Thr Leu Ile 1980	Ala Ala Met Ala 1885 Ile Asn His Ser Ser 1965	Ala Gly Ser 1870 Ile Leu Arg Tyr Ser 1950 Ser	Leu Tyr 1855 Gly Leu Arg Leu Val 1935 Leu Glu Trp	Asp 1840 Gly Glu Ser Arg Ile 1920 Pro Thr Cys
1810 Ala Ala Thr Ala 1825 Ser Val Gly Leu Ala Gly Val Ala 1860 Val Pro Ser Thr 1875 Pro Gly Ala Leu 1890 Arg Val Gly Pro 1905 Ala Phe Ala Ser Glu Ser Asp Ala 1940 Val Thr Gln Leu 1955 Thr Thr Pro Cys 1970 Cys Glu Val Leu	Phe Val 1830 Gly Lys 1845 Gly Ala Glu Asp Ala Val Gly Glu 1910 Arg Gly 1925 Ala Ala Leu Arg Ser Gly Ser Asp	1815 Gly Al Val Leu Va Leu Va 1895 Gly Al Asn Hi Arg Va Arg Le Ser Tr 1975 Phe Ly	la Gly eu Val al Ala 1865 al Asn 880 al Val la Val is Val al Thr 1945 eu His 960 rp Leu	Leu Ala 183 Asp Ile 1850 Phe Lys Leu Leu Phe Ala Gln Trp 1930 Ala Ile Gln Trp Arg Asp	1820 Gly 5 Leu Ile Pro Ser 1900 Met 5 Thr Leu Ile 1980 Lys	Ala Ala Met Ala 1885 Ile Asn His Ser Ser 1965	Ala Gly Ser 1870 Ile Leu Arg Tyr Ser 1950 Ser	Leu Tyr 1855 Gly Leu Arg Leu Val 1935 Leu Glu Trp	Asp 1840 Gly Glu Ser Arg Ile 1920 Pro Thr Cys Ile
Ala Ala Thr Ala 1825 Ser Val Gly Leu Ala Gly Val Ala 1860 Val Pro Ser Thr 1875 Pro Gly Ala Leu 1890 Arg Val Gly Pro 1905 Ala Phe Ala Ser Glu Ser Asp Ala 1940 Val Thr Gln Leu 1955 Thr Thr Pro Cys 1970 Cys Glu Val Leu 1985	Phe Val 1830 Gly Lys 1845 Gly Ala Glu Asp Ala Val Gly Glu 1910 Arg Gly 1925 Ala Ala Leu Arg Ser Gly 1990	1815 Gly Al Val Leu Va Leu Va 1895 Gly Al Asn Hi Arg Va Arg Le Ser Tr 1975 Phe Ly	la Gly eu Val al Ala 1865 al Asn 880 al Val la Val is Val al Thr 1945 eu His 960 rp Leu	Leu Ala 183 Asp Ile 1850 Phe Lys Leu Leu Phe Ala Gln Trp 1930 Ala Ile Gln Trp Arg Asp Trp Leu 199	1820 Gly 5 Leu Ile Pro Ser 1900 Met 5 Thr Leu Ile 1980 Lys	Ala Ala Met Ala 1885 Ile Asn His Ser 1965 Trp Ala	Ala Gly Ser 1870 Ile Leu Arg Tyr Ser 1950 Ser Asp Lys	Leu Tyr 1855 Gly Leu Arg Leu Val 1935 Leu Glu Trp Leu	Asp 1840 Gly Glu Ser Arg Ile 1920 Pro Thr Cys Ile Met 2000
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Pro Arg Thr Cys Lys Asn Met Trp Ser Gly Thr Phe Phe Ile Asn Ala
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                                           2060
Tyr Thr Thr Gly Pro Cys Thr Pro Leu Pro Ala Pro Asn Tyr Lys Phe
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                                       2075
Ala Leu Trp Arg Val Ser Ala Glu Glu Tyr Val Glu Ile Arg Arg Val
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Gly Asp Phe His Tyr Val Ser Gly Met Thr Thr Asp Asn Leu Lys Cys
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Pro Cys Gln Ile Pro Ser Pro Glu Phe Phe Thr Glu Leu Asp Gly Val
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Arg Leu His Arg Phe Ala Pro Pro Cys Lys Pro Leu Leu Arg Glu Glu
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Val Ser Phe Arg Val Gly Leu His Glu Tyr Pro Val Gly Ser Gln Leu
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Pro Cys Glu Pro Glu Pro Asp Val Ala Val Leu Thr Ser Met Leu Thr
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Asp Pro Ser His Ile Thr Ala Glu Ala Ala Gly Arg Arg Leu Ala Arg
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                               2185
Gly Ser Pro Pro Ser Met Ala Ser Ser Ser Ala Ser Gln Leu Ser Ala
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       2195
                                               2205
Pro Ser Leu Lys Ala Thr Cys Thr Ala Asn His Asp Ser Pro Asp Ala
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                                           2220
Glu Leu Ile Glu Ala Asn Leu Leu Trp Arg Gln Glu Met Gly Gly Asn
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                                       2235
Ile Thr Arg Val Glu Ser Glu Asn Lys Val Val Ile Leu Asp Ser Phe
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                                   2250
Asp Pro Leu Val Ala Glu Glu Asp Glu Arg Glu Val Ser Val Pro Ala
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                               2265
                                                   2270
Glu Ile Leu Arg Lys Ser Arg Arg Phe Ala Pro Ala Leu Pro Val Trp
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                                               2285
Ala Arg Pro Asp Tyr Asn Pro Leu Leu Val Glu Thr Trp Lys Lys Pro
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Asp Tyr Glu Pro Pro Val Val His Gly Cys Pro Leu Pro Pro Arg
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Ser Pro Pro Val Pro Pro Pro Arg Lys Lys Arg Thr Val Val Leu Thr
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Glu Ser Thr Leu Pro Thr Ala Leu Ala Glu Leu Ala Thr Lys Ser Phe
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Ser Glu Pro Ala Pro Ser Gly Cys Pro Pro Asp Ser Asp Val Glu Ser
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Tyr Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu
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Ser Asp Gly Ser Trp Ser Thr Val Ser Ser Gly Ala Asp Thr Glu Asp
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Val Val Cys Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Val Thr
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Pro Cys Ala Ala Glu Glu Gln Lys Leu Pro Ile Asn Ala Leu Ser Asn
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Ser Leu Leu Arg His His Asn Leu Val Tyr Ser Thr Thr Ser Arg Ser
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Ala Cys Gln Arg Lys Lys Lys Val Thr Phe Asp Arg Leu Gln Val Leu
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Lys Val Lys Ala 250		u Ser Val 2505		Cys Ser Let 2510	ı Ala
Pro Pro His Sec 2515	r Ala Lys Se	r Lys Phe (2520	Gly Tyr Gly	Ala Lys Asp 2525	Val
Arg Cys His Ala 2530	25	35	254	0	
Asp Leu Leu Glu 2545	2550		2555		2560
Lys Asn Glu Va	Phe Cys Va 2565		Glu Lys Gly 2570	Gly Arg Lys	
Ala Arg Leu Ile 258		o Asp Leu 2585		Val Cys Glu 2590	i Lys
Met Ala Leu Ty: 2595	Asp Val Va	l Ser Lys : 2600	Leu Pro Leu	Ala Val Met 2605	Gly
Ser Ser Tyr Gly 2610	Phe Gln Ty 26		Gly Gln Arg 262		e Leu
Val Gln Ala Trp 2625	Lys Ser Ly 2630	s Lys Thr	Pro Met Gly 2635	Leu Ser Ty	2640
Thr Arg Cys Pho	e Asp Ser Th		Glu Ser Asp 2650	Ile Arg Thi	
Glu Ala Ile Ty: 260		s Asp Leu . 2665	_	Ala Arg Val	l Ala
Ile Lys Ser Let 2675	Thr Glu Ar	g Leu Tyr 2680	Val Gly Gly	Pro Leu Thi 2685	Asn
Ser Arg Gly Glu 2690	Asn Cys Gl 26		Arg Cys Arg 270	_	y Val
Leu Thr Thr Set 2705	Cys Gly As: 2710	n Thr Leu	Thr Arg Tyr 2715	Ile Lys Ala	a Arg 2720
Ala Ala Cys Arg	2725		2730	273	35
Gly Asp Asp Let 27	10	2745		2750	
Ala Ala Ser Let 2755	_	2760		2765	
Pro Pro Gly Asp 2770	27	75	278	0	
Ser Cys Ser Se: 2785	Asn Val Se 2790	r Val Ala	His Asp Gly 2795	Ala Gly Lys	2800
Val Tyr Tyr Le	Thr Arg As 2805	_	Thr Pro Leu 2810	Ala Arg Ala 281	
Trp Glu Thr Ala 28:		r Pro Val . 2825		Leu Gly Asr 2830	ı Ile
Ile Met Phe Ala 2835	a Pro Thr Le	u Trp Ala . 2840	Arg Met Ile	Leu Met Thi 2845	His
Phe Phe Ser Val	Leu Ile Ala 28		Gln Leu Glu 286		ı Asn
Cys Glu Ile Ty: 2865	Gly Ala Cy 2870	s Tyr Ser	Ile Glu Pro 2875	Leu Asp Leu	2880
Pro Ile Ile Gli				Ser Leu His	
Tyr Ser Pro Gly		n Arg Val . 2905	_	Leu Arg Lys 2910	Leu
Gly Val Pro Pro 2915		a Trp Arg	His Arg Ala	Trp Ser Val	Arg

```
Ala Arg Leu Leu Ala Arg Gly Gly Lys Ala Ala Ile Cys Gly Lys Tyr
    2930
                        2935
Leu Phe Asn Trp Ala Val Arg Thr Lys Leu Lys Leu Thr Pro Ile Thr
                   2950
                                       2955
Ala Ala Gly Arg Leu Asp Leu Ser Gly Trp Phe Thr Ala Gly Tyr Ser
                                  2970
               2965
Gly Gly Asp Ile Tyr His Ser Val Ser His Ala Arg Pro Arg Trp Phe
           2980
                               2985
Trp Phe Cys Leu Leu Leu Ala Ala Gly Val Gly Ile Tyr Leu Leu
                            3000
Pro Asn Arg
    3010
<210> 2
<211> 182
<212> PRT
<213> Artificial Sequence
<223> Hepatitis C virus core protein sequence
<400> 2
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
                                    10
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
                       55
Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
                                        75
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
                                    90
Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
           100
                               105
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
                           120
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
                        135
Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
                   150
                                        155
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
               165
                                    170
Phe Leu Leu Ala Leu Leu
            180
<210> 3
<211> 197
<212> PRT
<213> Artificial Sequence
<223> Hepatitis C virus E1 protein sequence
```

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Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Gln Val Arg Asn Ser Ser
Gly Leu Tyr His Val Thr Asn Asp Cys Pro Asn Ser Ser Val Val Tyr
                                25
Glu Ala Ala Asp Ala Ile Leu His Thr Pro Gly Cys Val Pro Cys Val
                            40
Arg Glu Gly Asn Ala Ser Arg Cys Trp Val Ala Val Thr Pro Thr Val
                        55
Ala Thr Arg Asp Gly Lys Leu Pro Thr Thr Gln Leu Arg Arg His Ile
                    70
                                        75
Asp Leu Leu Val Gly Ser Ala Thr Leu Cys Ser Ala Leu Tyr Val Gly
                                    90
Asp Leu Cys Gly Ser Val Phe Leu Val Gly Gln Leu Phe Thr Phe Ser
                                105
Pro Arg His His Trp Thr Thr Gln Asp Cys Asn Cys Ser Ile Tyr Pro
                            120
Gly His Ile Thr Gly His Arg Met Ala Trp Asn Met Met Met Asn Trp
                        135
Ser Pro Thr Ala Ala Leu Val Val Ala Gln Leu Leu Arg Ile Pro Gln
                    150
                                        155
Ala Ile Met Asp Met Ile Ala Gly Ala His Trp Gly Val Leu Ala Gly
               165
                                    170
Ile Lys Tyr Phe Ser Met Val Gly Asn Trp Ala Lys Val Leu Val Val
            180
                                185
Leu Leu Phe Ala
        195
<210> 4
<211> 350
<212> PRT
<213> Artificial Sequence
<223> Hepatitis C virus E2 protein sequence
<400> 4
Gly Val Asp Ala Glu Thr His Val Thr Gly Gly Asn Ala Gly Arg Thr
Thr Ala Gly Leu Val Gly Leu Leu Thr Pro Gly Ala Lys Gln Asn Ile
Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn Ser Thr Ala Leu
                            40
Asn Cys Asn Glu Ser Leu Asn Thr Gly Trp Leu Ala Gly Leu Phe Tyr
Gln His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys
                    70
                                        75
Arg Arg Leu Thr Asp Phe Ala Gln Gly Trp Gly Pro Ile Ser Tyr Ala
                                    90
Asn Gly Ser Gly Leu Asp Glu Arg Pro Tyr Cys Trp His Tyr Pro Pro
                                105
Arg Pro Cys Gly Ile Val Pro Ala Lys Ser Val Cys Gly Pro Val Tyr
                            120
Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Ser Gly
Ala Pro Thr Tyr Ser Trp Gly Ala Asn Asp Thr Asp Val Phe Val Leu
```

<400> 3

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150
                                        155
Asn Asn Thr Arg Pro Pro Leu Gly Asn Trp Phe Gly Cys Thr Trp Met
                                    170
                165
Asn Ser Thr Gly Phe Thr Lys Val Cys Gly Ala Pro Pro Cys Val Ile
                                185
Gly Gly Val Gly Asn Asn Thr Leu Leu Cys Pro Thr Asp Cys Phe Arg
                            200
                                                205
Lys Tyr Pro Glu Ala Thr Tyr Ser Arg Cys Gly Ser Gly Pro Arg Ile
                        215
                                            220
Thr Pro Arg Cys Met Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro
                    230
                                        235
Cys Thr Ile Asn Tyr Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly
                245
                                    250
Val Glu His Arg Leu Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg
                                265
Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu
                            280
                                                285
Ser Thr Thr Gln Trp Gln Val Leu Pro Cys Ser Phe Thr Thr Leu Pro
                        295
Ala Leu Ser Thr Gly Leu Ile His Leu His Gln Asn Ile Val Asp Val
                    310
                                        315
Gln Tyr Leu Tyr Gly Val Gly Ser Ser Ile Ala Ser Trp Ala Ile Lys
               325
                                    330
Trp Glu Tyr Val Val Leu Leu Phe Leu Leu Leu Ala Asp Ala
                                345
```

<210> 5

<211> 315

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis C virus NS2 protein sequence

<400> 5

Arg Val Cys Ser Cys Leu Trp Met Met Leu Leu Ile Ser Gln Ala Glu 10 Ala Ala Leu Glu Asn Leu Val Ile Leu Asn Ala Ala Ser Leu Ala Gly 25 Thr His Gly Leu Val Ser Phe Leu Val Phe Phe Cys Phe Ala Trp Tyr Leu Lys Gly Arg Trp Val Pro Gly Ala Val Tyr Ala Leu Tyr Gly Met 55 Trp Pro Leu Leu Leu Leu Leu Ala Leu Pro Gln Arg Ala Tyr Ala 70 75 Leu Asp Thr Glu Val Ala Ala Ser Cys Gly Gly Val Val Leu Val Gly 90 Leu Met Ala Leu Thr Leu Ser Pro Tyr Tyr Lys Arg Tyr Ile Ser Trp 100 105 Cys Met Trp Trp Leu Gln Tyr Phe Leu Thr Arg Val Glu Ala Gln Leu 120 125 His Val Trp Val Pro Pro Leu Asn Val Arg Gly Gly Arg Asp Ala Val 135 140 Ile Leu Leu Thr Cys Val Val His Pro Ala Leu Val Phe Asp Ile Thr Lys Leu Leu Ala Ile Phe Gly Pro Leu Trp Ile Leu Gln Ala Ser

```
Leu Leu Lys Val Pro Tyr Phe Val Arg Val Gln Gly Leu Leu Arg Ile
                                185
            180
Cys Ala Leu Ala Arg Lys Ile Ala Gly Gly His Tyr Val Gln Met Ala
                            200
Ile Ile Lys Leu Gly Ala Leu Thr Gly Thr Cys Val Tyr Asn His Leu
                        215
                                            220
Ala Pro Leu Arg Asp Trp Ala His Asn Gly Leu Arg Asp Leu Ala Val
                    230
                                        235
Ala Val Glu Pro Val Val Phe Ser Arg Met Glu Thr Lys Leu Ile Thr
                                    250
                245
Trp Gly Ala Asp Thr Ala Ala Cys Gly Asp Ile Ile Asn Gly Leu Pro
                                265
            260
Val Ser Ala Arg Arg Gly Gln Glu Ile Leu Leu Gly Pro Ala Asp Gly
                            280
Met Val Ser Lys Gly Trp Arg Leu Leu Ala Pro Ile Thr Ala Tyr Ala
                                            300
                        295
Gln Gln Thr Arg Gly Leu Leu Gly Cys Ile Ile
                    310
<210> 6
<211> 613
<212> PRT
<213> Artificial Sequence
<223> Hepatitis C virus NS3 protein sequence
<400> 6
Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu Val Gln
                                    10
Ile Val Ser Thr Ala Thr Gln Thr Phe Leu Ala Thr Cys Ile Asn Gly
                                25
Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile Ala Ser
                            40
Pro Lys Gly Pro Val Ile Gln Thr Tyr Thr Asn Val Asp Gln Asp Leu
                        55
                                            60
Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro Cys Thr
                                        75
Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp Val Ile
                                    90
Pro Val Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser Pro Arg
                                105
Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu Cys Pro
                            120
Thr Gly His Ala Val Gly Leu Phe Arg Ala Ala Val Cys Thr Arg Gly
                        135
                                            140
Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Asn Leu Glu Thr Thr
                    150
                                        155
Met Arg Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala Val Pro
                165
                                    170
Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser Gly Lys
                                185
Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Lys Gly Tyr Lys Val Leu
                            200
Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala Tyr Met
```

170

```
215
                                            220
Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val Arg Thr
                   230
                                        235
Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys Phe Leu
               245
                                   250
Ala Asp Ala Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Cys Asp
                               265
Glu Cys His Ser Thr Asp Ala Thr Ser Ile Ser Gly Ile Gly Thr Val
                           280
Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala Thr
                       295
                                           300
Ala Thr Pro Pro Gly Ser Val Thr Val Ser His Pro Asn Ile Glu Glu
                   310
                                        315
Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys Ala Ile
                                    330
Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser
                                345
                                                    350
Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu Gly Ile
                           360
Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile Pro Thr
                       375
                                           380
Ser Gly Asp Val Val Val Ser Thr Asp Ala Leu Met Thr Gly Phe
                   390
                                       395
Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Cys Val Thr Gln
               405
                                   410
Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Thr
           420
                                425
Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr Gly
                           440
Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu Arg Pro
                       455
                                           460
Ser Gly Met Phe Asp Ser Ser Val Leu Cys Glu Cys Tyr Asp Ala Gly
                   470
                                       475
Cys Ala Trp Tyr Glu Leu Thr Pro Ala Glu Thr Thr Val Arg Leu Arg
               485
                                   490
Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu Gly
           500
                               505
Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile Asp Ala His Phe
                           520
Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Phe Pro Tyr Leu Val Ala
                       535
                                            540
Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser Trp
                   550
                                        555
Asp Gln Met Arg Lys Cys Leu Ile Arg Leu Lys Pro Thr Leu His Gly
               565
                                    570
Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn Glu Val Thr
                               585
Leu Thr His Pro Ile Thr Lys Tyr Ile Met Thr Cys Met Ser Ala Asp
                           600
                                               605
Leu Glu Val Val Thr
   610
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<210> 7

<211> 54

<212> PRT

<213> Artificial Sequence

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<220>
<223> Hepatitis C virus NS4A protein sequence
Ser Thr Trp Val Leu Val Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr
Cys Leu Ser Thr Gly Cys Val Val Ile Val Gly Arg Ile Val Leu Ser
                                25
Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu Val Leu Tyr Gln Glu Phe
                            40
Asp Glu Met Glu Glu Cys
    50
<210> 8
<211> 260
<212> PRT
<213> Artificial Sequence
<220>
<223> Hepatitis C virus NS4B protein sequence
<400> 8
Ser Gln His Leu Pro Tyr Ile Glu Gln Gly Met Met Leu Ala Glu Gln
Phe Lys Gln Lys Ala Leu Gly Leu Leu Gln Thr Ala Ser Arg His Ala
Glu Val Ile Thr Pro Ala Val Gln Thr Asn Trp Gln Lys Leu Glu Val
                            40
Phe Trp Ala Lys His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu
                        55
Ala Gly Leu Ser Thr Leu Pro Gly Asn Pro Ala Ile Ala Ser Leu Met
                    70
                                        75
Ala Phe Thr Ala Ala Val Thr Ser Pro Leu Thr Thr Gly Gln Thr Leu
                                    90
Leu Phe Asn Ile Leu Gly Gly Trp Val Ala Ala Gln Leu Ala Ala Pro
                                105
Gly Ala Ala Thr Ala Phe Val Gly Ala Gly Leu Ala Gly Ala Ala Leu
                            120
Asp Ser Val Gly Leu Gly Lys Val Leu Val Asp Ile Leu Ala Gly Tyr
                        135
                                            140
Gly Ala Gly Val Ala Gly Ala Leu Val Ala Phe Lys Ile Met Ser Gly
                    150
                                        155
Glu Val Pro Ser Thr Glu Asp Leu Val Asn Leu Leu Pro Ala Ile Leu
                                    170
                165
Ser Pro Gly Ala Leu Ala Val Gly Val Val Phe Ala Ser Ile Leu Arg
                                185
Arg Arg Val Gly Pro Gly Glu Gly Ala Val Gln Trp Met Asn Arg Leu
                            200
                                                205
Ile Ala Phe Ala Ser Arg Gly Asn His Val Ser Pro Thr His Tyr Val
                        215
Pro Glu Ser Asp Ala Ala Ala Arg Val Thr Ala Ile Leu Ser Ser Leu
                    230
                                        235
Thr Val Thr Gln Leu Leu Arg Arg Leu His Gln Trp Ile Ser Ser Glu
                                    250
```

Cys Thr Thr Pro

```
<210> 9
<211> 1040
<212> PRT
<213> Artificial Sequence
<223> Hepatitis C virus NS5A/B protein sequence
Cys Ser Gly Ser Trp Leu Arg Asp Ile Trp Asp Trp Ile Cys Glu Val
Leu Ser Asp Phe Lys Thr Trp Leu Lys Ala Lys Leu Met Pro Gln Leu
Pro Gly Ile Pro Phe Val Ser Cys Gln Arg Gly Tyr Arg Gly Val Trp
                            40
Arg Gly Asp Gly Ile Met His Thr Arg Cys His Cys Gly Ala Glu Ile
                        55
Thr Gly His Val Lys Asn Gly Thr Met Arg Ile Val Gly Pro Arg Thr
                    70
Cys Lys Asn Met Trp Ser Gly Thr Phe Phe Ile Asn Ala Tyr Thr Thr
                                    90
                85
Gly Pro Cys Thr Pro Leu Pro Ala Pro Asn Tyr Lys Phe Ala Leu Trp
            100
                                105
Arg Val Ser Ala Glu Glu Tyr Val Glu Ile Arg Arg Val Gly Asp Phe
                            120
His Tyr Val Ser Gly Met Thr Thr Asp Asn Leu Lys Cys Pro Cys Gln
                        135
                                            140
Ile Pro Ser Pro Glu Phe Phe Thr Glu Leu Asp Gly Val Arg Leu His
                    150
                                        155
Arg Phe Ala Pro Pro Cys Lys Pro Leu Leu Arg Glu Glu Val Ser Phe
                165
                                    170
Arg Val Gly Leu His Glu Tyr Pro Val Gly Ser Gln Leu Pro Cys Glu
            180
                                185
                                                     190
Pro Glu Pro Asp Val Ala Val Leu Thr Ser Met Leu Thr Asp Pro Ser
                            200
                                                 205
His Ile Thr Ala Glu Ala Ala Gly Arg Arg Leu Ala Arg Gly Ser Pro
                        215
                                            220
Pro Ser Met Ala Ser Ser Ser Ala Ser Gln Leu Ser Ala Pro Ser Leu
                    230
                                         235
Lys Ala Thr Cys Thr Ala Asn His Asp Ser Pro Asp Ala Glu Leu Ile
                245
                                    250
Glu Ala Asn Leu Leu Trp Arg Gln Glu Met Gly Gly Asn Ile Thr Arg
                                265
Val Glu Ser Glu Asn Lys Val Val Ile Leu Asp Ser Phe Asp Pro Leu
                            280
                                                285
Val Ala Glu Glu Asp Glu Arg Glu Val Ser Val Pro Ala Glu Ile Leu
                        295
                                            300
Arg Lys Ser Arg Arg Phe Ala Pro Ala Leu Pro Val Trp Ala Arg Pro
                    310
                                        315
Asp Tyr Asn Pro Leu Leu Val Glu Thr Trp Lys Lys Pro Asp Tyr Glu
                                    330
                325
Pro Pro Val Val His Gly Cys Pro Leu Pro Pro Pro Arg Ser Pro Pro
                                345
Val Pro Pro Pro Arg Lys Lys Arg Thr Val Val Leu Thr Glu Ser Thr
```

```
355
                            360
Leu Pro Thr Ala Leu Ala Glu Leu Ala Thr Lys Ser Phe Gly Ser Ser
                        375
                                            380
Ser Thr Ser Gly Ile Thr Gly Asp Asn Thr Thr Ser Ser Glu Pro
                    390
                                        395
Ala Pro Ser Gly Cys Pro Pro Asp Ser Asp Val Glu Ser Tyr Ser Ser
                405
                                    410
Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu Ser Asp Gly
                                425
            420
Ser Trp Ser Thr Val Ser Ser Gly Ala Asp Thr Glu Asp Val Val Cys
                            440
Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Val Thr Pro Cys Ala
                        455
                                            460
Ala Glu Glu Gln Lys Leu Pro Ile Asn Ala Leu Ser Asn Ser Leu Leu
                    470
                                        475
Arg His His Asn Leu Val Tyr Ser Thr Thr Ser Arg Ser Ala Cys Gln
                                    490
Arg Lys Lys Val Thr Phe Asp Arg Leu Gln Val Leu Asp Ser His
           500
                                505
Tyr Gln Asp Val Leu Lys Glu Val Lys Ala Ala Ala Ser Lys Val Lys
                            520
Ala Asn Leu Leu Ser Val Glu Glu Ala Cys Ser Leu Ala Pro Pro His
                        535
                                            540
Ser Ala Lys Ser Lys Phe Gly Tyr Gly Ala Lys Asp Val Arg Cys His
                    550
                                        555
Ala Arg Lys Ala Val Ala His Ile Asn Ser Val Trp Lys Asp Leu Leu
                                    570
Glu Asp Ser Val Thr Pro Ile Asp Thr Thr Ile Met Ala Lys Asn Glu
                                585
Val Phe Cys Val Gln Pro Glu Lys Gly Gly Arg Lys Pro Ala Arg Leu
                            600
Ile Val Phe Pro Asp Leu Gly Val Arg Val Cys Glu Lys Met Ala Leu
                       615
                                            620
Tyr Asp Val Val Ser Lys Leu Pro Leu Ala Val Met Gly Ser Ser Tyr
                    630
                                        635
Gly Phe Gln Tyr Ser Pro Gly Gln Arg Val Glu Phe Leu Val Gln Ala
                645
                                    650
Trp Lys Ser Lys Lys Thr Pro Met Gly Leu Ser Tyr Asp Thr Arg Cys
            660
                                665
Phe Asp Ser Thr Val Thr Glu Ser Asp Ile Arg Thr Glu Glu Ala Ile
                            680
Tyr Gln Cys Cys Asp Leu Asp Pro Gln Ala Arg Val Ala Ile Lys Ser
                                            700
                        695
Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Leu Thr Asn Ser Arg Gly
                    710
                                        715
Glu Asn Cys Gly Tyr Arg Arg Cys Arg Ala Ser Arg Val Leu Thr Thr
                725
                                    730
Ser Cys Gly Asn Thr Leu Thr Arg Tyr Ile Lys Ala Arg Ala Ala Cys
            740
                                745
Arg Ala Ala Gly Leu Gln Asp Cys Thr Met Leu Val Cys Gly Asp Asp
                            760
Leu Val Val Ile Cys Glu Ser Ala Gly Val Gln Glu Asp Ala Ala Ser
                        775
                                            780
Leu Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser Ala Pro Pro Gly
                    790
                                        795
Asp Pro Pro Gln Pro Glu Tyr Asp Leu Glu Leu Ile Thr Ser Cys Ser
                805
                                    810
```

```
825
Leu Thr Arg Asp Pro Thr Thr Pro Leu Ala Arg Ala Ala Trp Glu Thr
                            840
Ala Arq His Thr Pro Val Asn Ser Trp Leu Gly Asn Ile Ile Met Phe
                        855
Ala Pro Thr Leu Trp Ala Arg Met Ile Leu Met Thr His Phe Phe Ser
                    870
                                        875
Val Leu Ile Ala Arg Asp Gln Leu Glu Gln Ala Leu Asn Cys Glu Ile
                885
                                    890
Tyr Gly Ala Cys Tyr Ser Ile Glu Pro Leu Asp Leu Pro Pro Ile Ile
                                905
Gln Arg Leu His Gly Leu Ser Ala Phe Ser Leu His Ser Tyr Ser Pro
                            920
Gly Glu Ile Asn Arg Val Ala Ala Cys Leu Arg Lys Leu Gly Val Pro
                        935
                                            940
Pro Leu Arg Ala Trp Arg His Arg Ala Trp Ser Val Arg Ala Arg Leu
                    950
                                        955
Leu Ala Arg Gly Gly Lys Ala Ala Ile Cys Gly Lys Tyr Leu Phe Asn
               965
                                    970
Trp Ala Val Arg Thr Lys Leu Lys Leu Thr Pro Ile Thr Ala Ala Gly
            980
                                985
                                                    990
Arg Leu Asp Leu Ser Gly Trp Phe Thr Ala Gly Tyr Ser Gly Gly Asp
                            1000
                                                1005
Ile Tyr His Ser Val Ser His Ala Arg Pro Arg Trp Phe Trp Phe Cys
                        1015
                                            1020
Leu Leu Leu Ala Ala Gly Val Gly Ile Tyr Leu Leu Pro Asn Arg
                    1030
                                        1035
<210> 10
<211> 226
<212> PRT
<213> Artificial Sequence
<220>
<223> Hepatitis B virus S antigen (HBsAg) sequence
<400> 10
Met Glu Asn Ile Thr Ser Gly Phe Leu Gly Pro Leu Leu Val Leu Gln
Ala Gly Phe Phe Leu Leu Thr Arg Ile Leu Thr Ile Pro Gln Ser Leu
                                25
Asp Ser Trp Trp Thr Ser Leu Asn Phe Leu Gly Gly Thr Thr Val Cys
                            40
Leu Gly Gln Asn Ser Gln Ser Pro Thr Ser Asn His Ser Pro Thr Ser
                        55
Cys Pro Pro Thr Cys Pro Gly Tyr Arg Trp Met Cys Leu Arg Arg Phe
                    70
                                        75
Ile Ile Phe Leu Phe Ile Leu Leu Cys Leu Ile Phe Leu Leu Val
                85
                                    90
Leu Leu Asp Tyr Gln Gly Met Leu Pro Val Cys Pro Leu Ile Pro Gly
                                105
Ser Ser Thr Thr Ser Thr Gly Pro Cys Arg Thr Cys Met Thr Thr Ala
                            120
```

Ser Asn Val Ser Val Ala His Asp Gly Ala Gly Lys Arg Val Tyr Tyr

Gln Gly Thr Ser Met Tyr Pro Ser Cys Cys Cys Thr Lys Pro Ser Asp

135

```
Gly Asn Cys Thr Cys Ile Pro Ile Pro Ser Ser Trp Ala Phe Gly Lys
                                       155
                    150
Phe Leu Trp Glu Trp Ala Ser Ala Arg Phe Ser Trp Leu Ser Leu Leu
                                   170
Val Pro Phe Val Gln Trp Phe Val Gly Leu Ser Pro Thr Val Trp Leu
                                185
Ser Val Ile Trp Met Met Trp Tyr Trp Gly Pro Ser Leu Tyr Ser Ile
                           200
Leu Ser Pro Phe Leu Pro Leu Pro Ile Phe Phe Cys Leu Trp Val
                       215
                                            220
Tyr Ile
225
<210> 11
<211> 212
<212> PRT
<213> Artificial Sequence
<223> Hepatitis B virus C antigen and e antigen
      (HBcAg/HBeAg) sequence
Met Gln Leu Phe His Leu Cys Leu Ile Ile Ser Cys Ser Cys Pro Thr
Val Gln Ala Ser Lys Leu Cys Leu Gly Trp Leu Trp Gly Met Asp Ile
Asp Pro Tyr Lys Glu Phe Gly Ala Thr Val Glu Leu Leu Ser Phe Leu
                            40
Pro Ser Asp Phe Phe Pro Ser Val Arg Asp Leu Leu Asp Thr Ala Ser
                       55
Ala Leu Tyr Arg Glu Ala Leu Glu Ser Pro Glu His Cys Ser Pro His
His Thr Ala Leu Arg Gln Ala Ile Leu Cys Trp Gly Glu Leu Met Thr
                85
                                    90
Leu Ala Thr Trp Val Gly Val Asn Leu Glu Asp Pro Ala Ser Arg Asp
Leu Val Val Ser Tyr Val Asn Thr Asn Met Gly Leu Lys Phe Arg Gln
                            120
Leu Leu Trp Phe His Ile Ser Cys Leu Thr Phe Gly Arg Glu Thr Val
                        135
                                            140
Ile Glu Tyr Leu Val Ser Phe Gly Val Trp Ile Arg Thr Pro Pro Ala
                   150
                                       155
Tyr Arg Pro Pro Asn Ala Pro Ile Leu Ser Thr Leu Pro Glu Thr Thr
                                   170
               165
Val Val Arg Arg Gly Arg Ser Pro Arg Arg Thr Pro Ser Pro
                               185
Arg Arg Arg Ser Gln Ser Pro Arg Arg Arg Ser Gln Ser Arg
                            200
                                                205
       195
Glu Ser Gln Cys
    210
<210> 12
<211> 2227
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<212> PRT

<220>

<223> Hepatitis A virus sequence

<400> 12 Met Asn Met Ser Lys Gln Gly Ile Phe Gln Thr Val Gly Ser Gly Leu 10 Asp His Ile Leu Ser Leu Ala Asp Ile Glu Glu Glu Gln Met Ile Gln 25 Ser Val Asp Arg Thr Ala Val Thr Gly Ala Ser Tyr Phe Thr Ser Val Asp Gln Ser Ser Val His Thr Ala Glu Val Gly Ser His Gln Ile Glu 55 Pro Leu Lys Thr Ser Val Asp Lys Pro Gly Ser Lys Lys Thr Gln Gly 70 75 Glu Lys Phe Phe Leu Ile His Ser Ala Asp Trp Leu Thr Thr His Ala 90 Leu Phe His Glu Val Ala Lys Leu Asp Val Val Lys Leu Leu Tyr Asn 105 100 Glu Gln Phe Ala Val Gln Gly Leu Leu Arg Tyr His Thr Tyr Ala Arg 120 Phe Gly Ile Glu Ile Gln Val Gln Ile Asn Pro Thr Pro Phe Gln Gln 135 140 Gly Gly Leu Ile Cys Ala Met Val Pro Gly Asp Gln Ser Tyr Gly Ser 150 155 Ile Ala Ser Leu Thr Val Tyr Pro His Gly Leu Leu Asn Cys Asn Ile 170 Asn Asn Val Val Arg Ile Lys Val Pro Phe Ile Tyr Thr Arg Gly Ala 180 185 Tyr His Phe Lys Asp Pro Gln Tyr Pro Val Trp Glu Leu Thr Ile Arg 200 Val Trp Ser Glu Leu Asn Ile Gly Thr Gly Thr Ser Ala Tyr Thr Ser 215 220 Leu Asn Val Leu Ala Arg Phe Thr Asp Leu Glu Leu His Gly Leu Thr 230 235 Pro Leu Ser Thr Gln Met Met Arg Asn Glu Phe Arg Val Ser Thr Thr 245 250 Glu Asn Val Val Asn Leu Ser Asn Tyr Glu Asp Ala Arg Ala Lys Met 260 265 Ser Phe Ala Leu Asp Gln Glu Asp Trp Lys Ser Asp Pro Ser Gln Gly 280 Gly Gly Ile Lys Ile Thr His Phe Thr Thr Trp Thr Ser Ile Pro Thr 295 300 Leu Ala Ala Gln Phe Pro Phe Asn Ala Ser Asp Ser Val Gly Gln Gln 310 315 Ile Lys Val Ile Pro Val Asp Pro Tyr Phe Phe Gln Met Thr Asn Thr 325 330 Asn Pro Asp Gln Lys Cys Ile Thr Ala Leu Ala Ser Ile Cys Gln Met 340 345 Phe Cys Phe Trp Arg Gly Asp Leu Val Phe Asp Phe Gln Val Phe Pro 360 Thr Lys Tyr His Ser Gly Arg Leu Leu Phe Cys Phe Val Pro Gly Asn 375 Glu Leu Ile Asp Val Thr Gly Ile Thr Leu Lys Gln Ala Thr Thr Ala 390 395 Pro Cys Ala Val Met Asp Ile Thr Gly Val Gln Ser Thr Leu Arg Phe

```
405
                                    410
Arq Val Pro Trp Ile Ser Asp Thr Pro Tyr Arg Val Asn Arg Tyr Thr
            420
                                425
Lys Ser Ala His Gln Lys Gly Glu Tyr Thr Ala Ile Gly Lys Leu Ile
                            440
Val Tyr Cys Tyr Asn Arg Leu Thr Ser Pro Ser Asn Val Ala Ser His
                        455
                                            460
Val Arg Val Asn Val Tyr Leu Ser Ala Ile Asn Leu Glu Cys Phe Ala
                    470
                                        475
Pro Leu Tyr His Ala Met Asp Val Thr Thr Gln Val Gly Asp Asp Ser
                                    490
                485
Gly Gly Phe Ser Thr Thr Val Ser Thr Glu Gln Asn Val Pro Asp Pro
                                505
Gln Val Gly Ile Thr Thr Met Arg Asp Leu Lys Gly Lys Ala Asn Arg
                            520
Gly Lys Met Asp Val Ser Gly Val Gln Ala Pro Arg Gly Ser Tyr Gln
                        535
Gln Gln Leu Asn Asp Pro Val Leu Ala Lys Lys Val Pro Glu Thr Phe
                    550
                                        555
Pro Glu Leu Lys Pro Gly Glu Ser Arg His Thr Ser Asp His Met Ser
                565
                                    570
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                                                                  7478
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